

Identification with X'Pert HighScore: Search-match and Identification Background Information

PANalytical

QUB XRD Course

Data Interpretation

Basic XRD Course 1

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Pattern Treatments - Data Reduction Functions

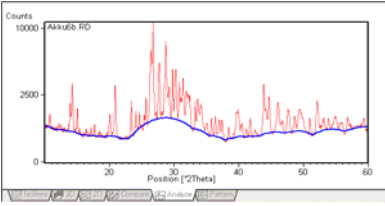
- Data reduction functions
 - Background determination
 - Peak search
 - Profile fitting
 - $K\alpha_2$ stripping
- Correction functions
 - Divergence slit
 -

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Background Determination

- Iterative approximation
- By peak search
- Manual



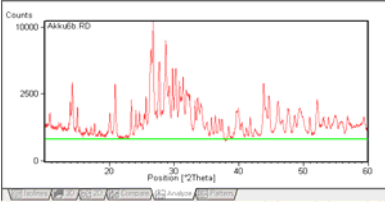
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Background Determination

- Iterative approximation
- **By peak search**
- Manual

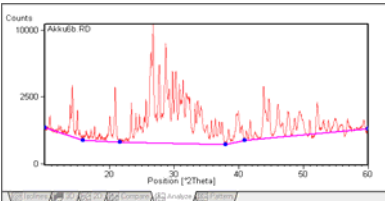


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Background Determination

- Iterative approximation
- By peak search
- **Manual**



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PANalytical

Pattern Treatments

- Data reduction functions
 - Background determination
 - **Peak search**
 - Profile fitting
 - $K\alpha_2$ stripping
 - Smoothing
- Correction functions
 - Divergence slit
 -
- Edit data

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Identification with X'Pert HighScore: Search-match and Identification Background Information

Peak Search Parameters

Search Peaks - [Default]

Minimum significance: 1.00 Search Peaks

Minimum tip width [°2Th.]: 0.01 Replace

Maximum tip width [°2Th.]: 1.00

Peak base width [°2Th.]: 2.00

Method: Minimum 2nd derivative Close

Tjral: More >>

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Peak Search - Purpose

- Find the **position**, estimated **width** and **height** of the peaks in your scan
- Input for
 - Phase Identification
 - Indexing (Unit cell determination)


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Peak Search - Algorithm

- The minimum of the second derivative curve gives the peak position
- This method removes the effect of sloping backgrounds

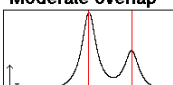
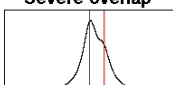
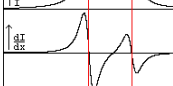
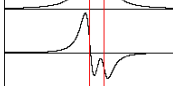
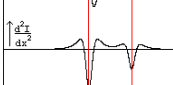
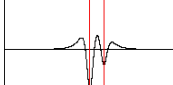
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Peak Search - Algorithm

- Overlapping peaks can be found with the 2nd derivative

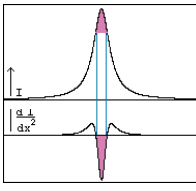
	Moderate overlap	Severe overlap
Intensity		
1 st derivative $\frac{dI}{dx}$		
2 nd derivative $\frac{d^2I}{dx^2}$		

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Peak Search - Significance

- Significance of the peak**
 - Area of 2nd derivative
 - Area of peak tip
- Peak tip width:**
 - Distance between points of inflection

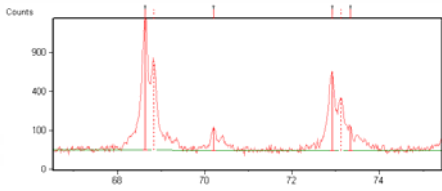


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Peak Search - Peak Derivation

After peak search X'Pert HighScore tries to identify all the peaks as $K\alpha_1$ or $K\alpha_2$ or $K\alpha_{mixed}$ (peak derivation)




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Pattern Treatments

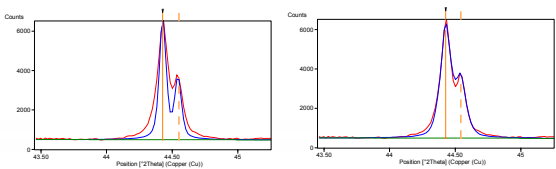
- Data reduction functions
 - Background determination
 - Peak search
 - Profile fitting
 - $K\alpha_2$ stripping
- Correction functions
 - Divergence slit
 -
- Edit data

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Profile fitting

- Obtain better peak parameters
 - Position
 - Height
 - Width (FWHM)



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Pattern Treatments

- Data reduction functions
 - Background determination
 - Peak search
 - Profile fitting
 - $K\alpha_2$ stripping
- Correction functions
 - Divergence slit
 -
- Edit data

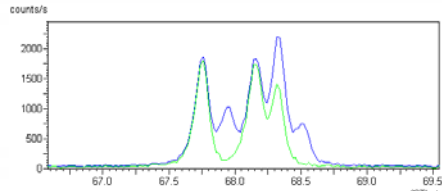
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$K\alpha_2$ Stripping - Purpose

- To remove the $K\alpha_2$ contribution from a scan
- To improve resolution

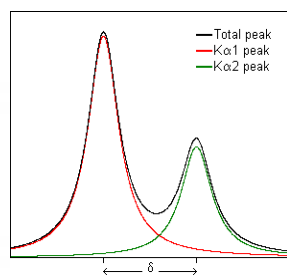


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$K\alpha_2$ Stripping - Algorithm

- Assume that the shape and width of the $K\alpha_1$ and $K\alpha_2$ peak are the same
- Use the known wavelengths to calculate the $K\alpha_2$ position from the $K\alpha_1$ position
- **Rachinger** method

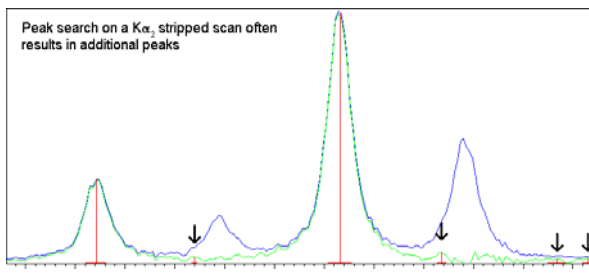


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$K\alpha_2$ Stripping - Example

Peak search on a $K\alpha_2$ stripped scan often results in additional peaks



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Pattern Treatments

- Data reduction functions
 - Background determination
 - Peak search
 - Profile fitting
 - $K\alpha_2$ stripping
- Correction functions
 - Divergence slit
 -
- Edit data


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Pattern Treatments - Scan Data Corrections

- Data reduction functions
- Correction functions
 - Divergence slit conversion
 - Step size interpolation
 - Sample displacement correction
 - Systematic error correction
 - Outlier correction
 - Temperature/pressure correction
 - Beam overflow correction

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Phase Identification

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Phase Identification - The Challenge

Find which phases from the reference database are present in your sample.

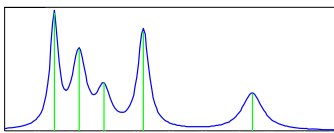
- Huge amount of patterns in the reference database
- Measuring errors
- Mixtures

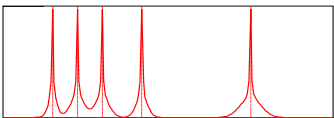
Features:	PDF-4+ 2008 with DDView+	PDF-2 Release 2008 with DDView
Total entries	285,402	211,107
Inorganic entries	256,934	182,634
Organic entries	32,408	32,189
Entries with atomic coordinates	114,630	0

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Finding Candidate Phases

Peak list 

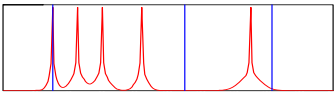
Probability curve of peak & profile data 

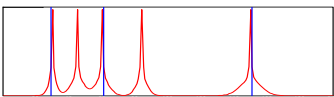
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Finding Candidate Phases - 3


Peak lists compared to probability curve

Low probability 

High probability 

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Semi-quantitative Analysis

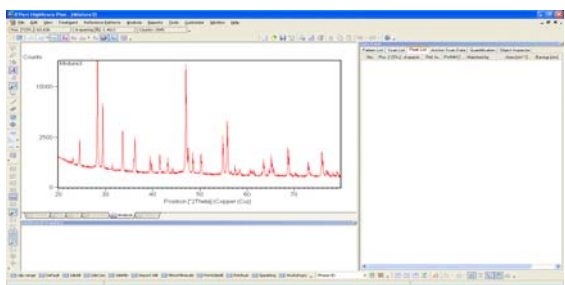
- Estimates the mass fractions of the identified phases
 - All phases must have been identified
 - Amorphous material is not taken into account
- The calculation is based on
 - Scale factors
 - Reference Intensity Ratio values

The RIR values are based on the relative net peak height ratio of the strongest line of the phase and of the strongest line of Corundum measured under the same conditions.

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Data ready for interpretation – X'Pert HighScore



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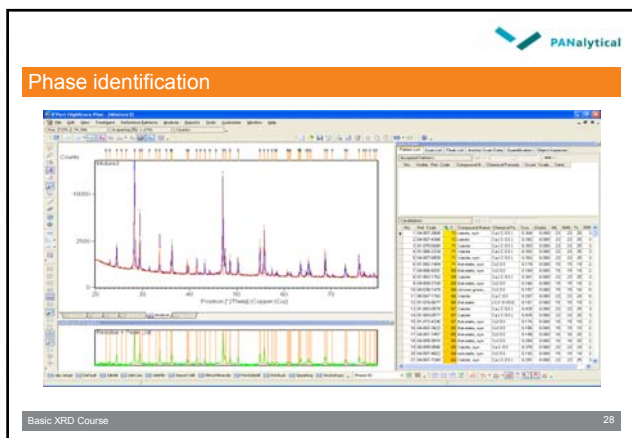


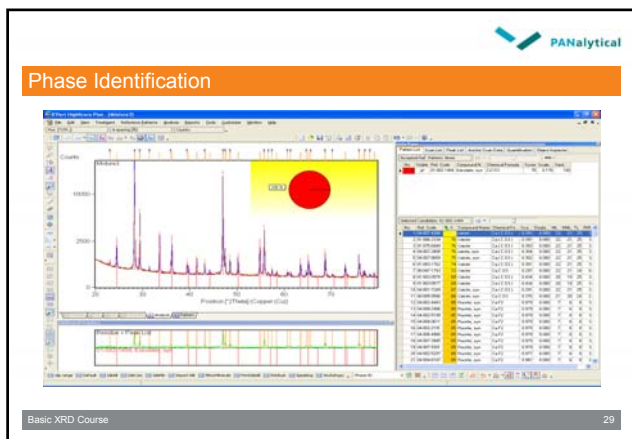
Pattern Treatment

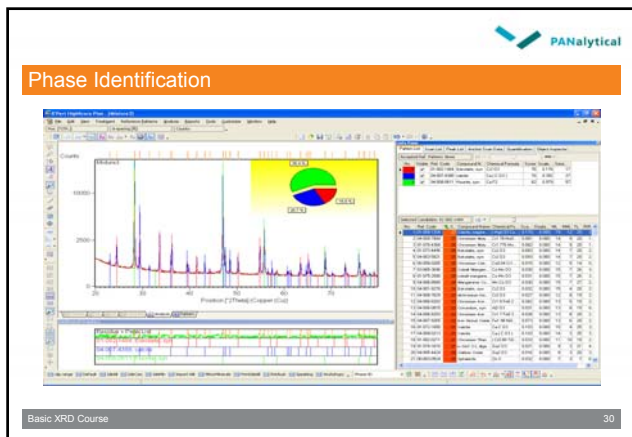


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Identification with X'Pert HighScore: Search-match and Identification Background Information







Identification with X'Pert HighScore: Search-match and Identification Background Information

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Rietveld


Introduction to Rietveld refinement

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Rietveld Refinement

- Devised by Hugo Rietveld, 1969
- Originally for Neutron diffraction
- Whole pattern structure refinement
- No crystallographic information is lost



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Rietveld refinement

- A calculated diffraction pattern is compared to measured data.
- The model used for the calculated pattern is adjusted to improve the fit between the two.

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Identification with X'Pert HighScore: Search-match and Identification Background Information

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What can Rietveld tell me?

- There are two ways that Rietveld can be used

Rietveld

Crystal Structure Refinement

Standardless Quantitative analysis

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Minimisation of differences

- The term that is minimised is S_y

$$S_y = \sum_i w_i (y_i - y_{cl})^2$$

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Quantitative methods - bulk samples

- Reference Intensity Ratio (RIR)
- Internal Standard method
- Straight line method
- Addition method
- Matrix flushing method
- General method


}

single line methods

- Rietveld method

← **Whole pattern method**


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Data collection

- High resolution
- High Intensity (> 10 000 counts)
- Large 2θ range (5-100°)
- At least 5-10 steps across FWHM
- Avoid preferred orientation
- Particle size 1-10µm

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Particle size

Intensity Measurements on Different Size Fractions of <325-Mesh Quartz Powder (after Klug and Alexander [1974], p. 366)

Specimen No.	15-50µ Fraction	5-50µ Fraction	5- 15µ Fraction	<5µ Fraction
1	7.612	8.688	10.841	11.055
2	8.373	9.040	11.336	11.040
3	8.265	10.232	11.046	11.386
4	9.333	9.333	11.597	11.212
5	4.823	8.530	11.541	11.460
6	11.123	8.617	11.336	11.260
7	11.051	11.598	11.686	11.241
8	5.773	7.818	11.288	11.428
9	8.527	8.021	11.126	11.406
10	10.255	10.190	10.878	11.444
Mean area	8.513	9.227	11.268	11.293
Mean deviation	1.545	0.929	0.236	0.132
Mean % deviation	18.2	10.1	2.1	1.2

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Calculated intensity

$$y_{ci} = s \sum_K L_K |F_K|^2 \phi(2\theta_i - 2\theta_K) P_K A + y_{bi}$$

S = Scale factor
 K represents the Miller indices
 L_K = Lorenz polarisation and multiplicity factors
 φ is the reflection profile function
 P_K is the preferred orientation function
 A is the absorption factor
 F_K is the structure factor of the Kth Bragg reflection
 Y_{bi} is the background intensity of the ith step

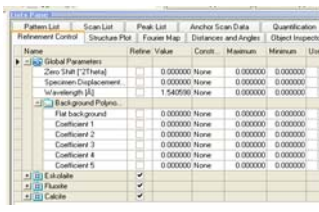
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Identification with X'Pert HighScore: Search-match and Identification Background Information




Refinement Parameters

- Global Parameters
 - Background
 - Zero shift
 - Specimen displacement

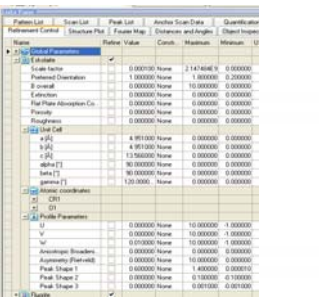


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Refinement Parameters

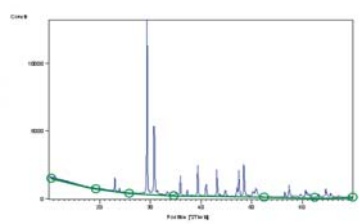
- Phase Related Parameters



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Manual Background



Background is set at the beginning of the refinement.

Straight line is drawn between set points.

Linear Interpolation

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Refined background

- Polynomial
- Chebyshev
- Shifted Chebyshev
- Amorphous sinc function
- Damped amorphous sinc function

All are made up of a series of coefficients


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Refined Background

$$I = A + B\theta + C\theta^2 + D\theta^3 + E\theta^4 \dots$$

1st coefficient (A) is referred to as the flat background parameter



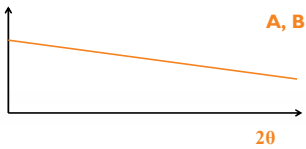
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PANalytical

Refined Background

$$I = A + B\theta + C\theta^2 + D\theta^3 + E\theta^4 \dots$$

1st coefficient (A) is referred to as the flat background parameter



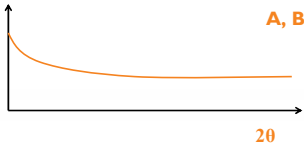
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Refined Background

$$I = A + B\theta + C\theta^2 + D\theta^3 + E\theta^4 \dots$$

1st coefficient (A) is referred to as the flat background parameter



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Manual Vs Refined

- Manual
 - Time consuming
 - Can miss weak reflections
 - Handles difficult backgrounds
- Refined
 - Can be automated
 - Adjustable throughout refinement

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Space Group


- Determines the symmetry of the structure
- Fixed at the start of the refinement
- 230 Space Groups

NaCl Fm3m Rutile P 4₂/mnm

Gypsum C 2/c

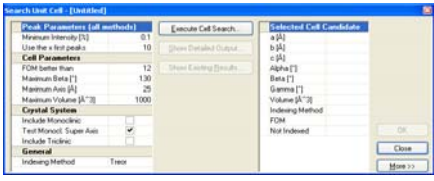
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Indexing

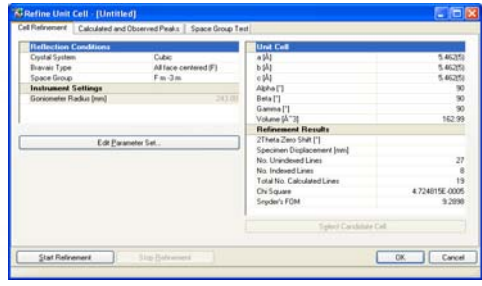
- Fitting peak list to unit cell dimensions
- Try different programs and compare results




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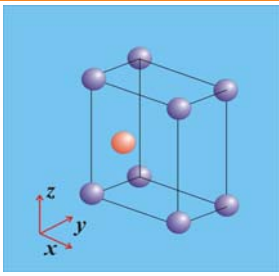
Cell Refinement



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Atomic Positions




Each atom in the structure must have its atomic coordinates recorded.

E.g.

- Special positions:**
 - (0, 0, 0)
 - ($\frac{1}{2}$, $\frac{1}{2}$, 0)
 - ($\frac{1}{3}$, $\frac{2}{3}$, 0.492)
- Non special positions:**
 - (0.2346, 0.987, 0.456)

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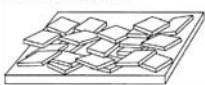
Identification with X'Pert HighScore: Search-match and Identification Background Information



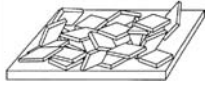
Preferred orientation

- Results from crystallites lining up with each other.
- May result from crystal growth mechanism or special cleavage planes.
- Use back-loading techniques and spin sample.
- Observe symmetry related peaks all high / low in intensity


Preferred orientation



Random orientation




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Peak Shape functions

- Gaussian (G)
- Lorentzian (L)
- Voigt
- Pseudo-Voigt
 - Mixture of Gaussian and Lorentzian
 - $pV = \eta L + (1-\eta)G$
- Pearson VII
- Mod-TCH pV

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Scale Factor

- Simple scalar variable
- Leads to quantification.

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Identification with X'Pert HighScore: Search-match and Identification Background Information

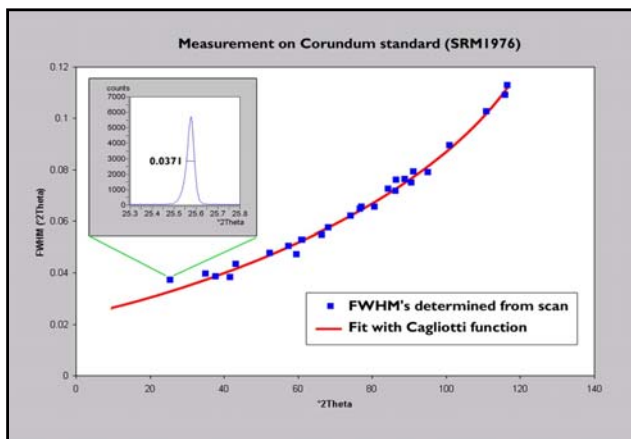



Peak Width

- Full Width Half Maximum (FWHM)
- FWHM is dependent on θ
- Cagliotti function

$$H^2 = U \tan^2\theta + V \tan\theta + W$$
- Often sufficient just to refine W


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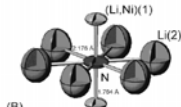


Temperature Factors

- Thermal vibration of the atoms
- Can be model Isotropic or Anisotropic
- Labelled U or B



No vibration **With vibration**



(B)

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Site Occupancy Factors SOF

- Is each site fully filled?
- Do two atoms share one site?
- Solid solutions

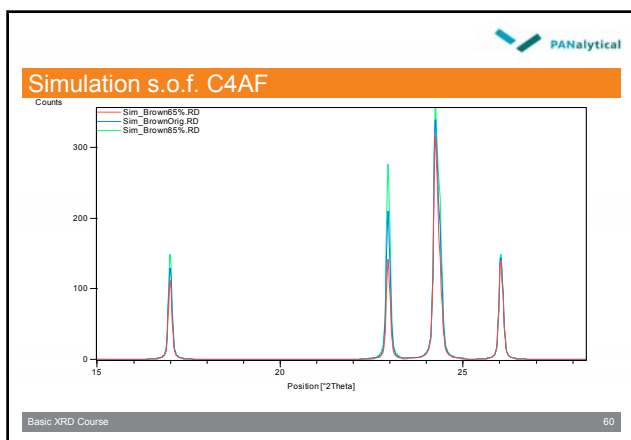
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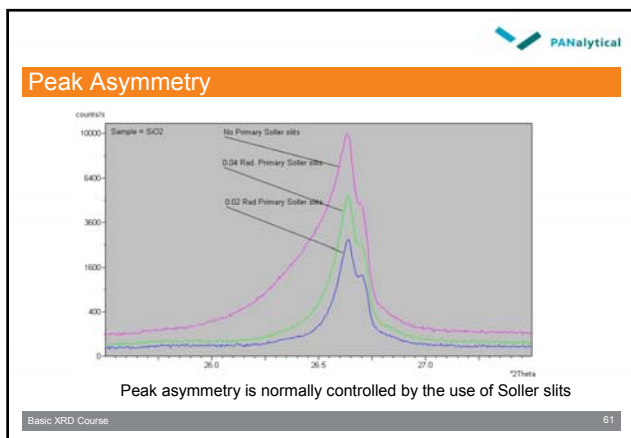
Site Occupancy Factor

- Brownmillerite Ca_4AF
- $4\text{CaO} \cdot \text{Al}_2\text{O}_3 \cdot \text{Fe}_2\text{O}_3$
- Iron and Aluminium share the same site
- $\text{Fe}^{3+} \leftrightarrow \text{Al}^{3+}$
- Need to link the occupancy together

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Identification with X'Pert HighScore: Search-match and Identification Background Information



-
- ### Refinement Strategy
- Start with few parameters
 - Increase number as refinement proceeds
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-
- ### Refinement Order / Strategy
1. Background + Scale factors
 2. Cell Parameters + Zero shift
 3. Peak width W
 4. Preferred orientation
 5. SOF
 6. Temperature Factors
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Assessing the quality of the refinement

- Four R-factors provide a measure of the fit
- R_p Profile factor
- R_{wp} Weighted profile factor
- R_B Bragg factor
- R_I Intensity factor


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Which factors are important?

- R_{wp} gives best indication of fit as the numerator is the residual being minimised during refinement.
- R_B and R_I are dependant only on the Bragg reflections so heavily biased towards the structural model.

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Why use Rietveld Quantification?

- When preferred orientation is unavoidable
- When standards are unavailable
- When occupancies vary
- When peaks overlap
- Data collection can now be done much quicker

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Identification with X'Pert HighScore: Search-match and Identification Background Information

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Potential problems

- Needs structure data
- Incorrect solutions can look good
- Can be numerically unstable

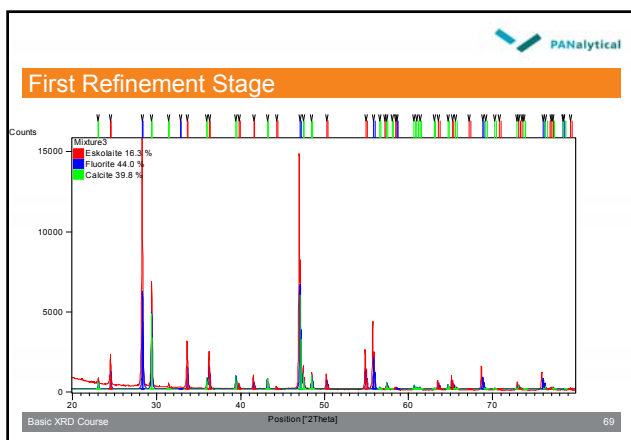
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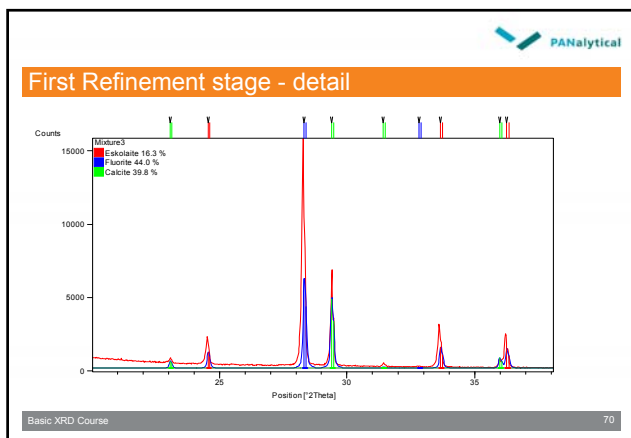
Unstable Refinements

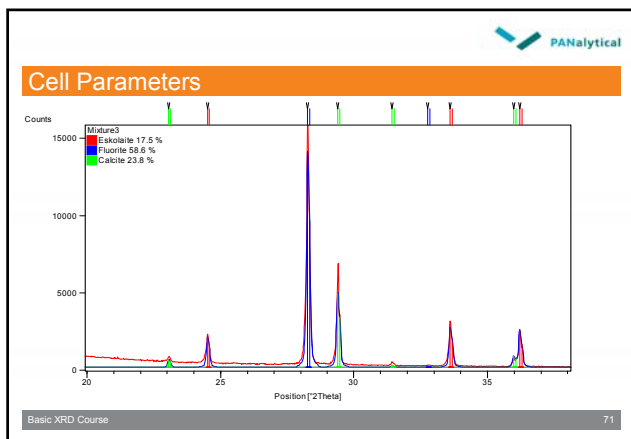
- Use variable limits
 - Compatible with automation
- Refine parameters individually then fix
- Change parameters manually and check by eye
- Use undo!

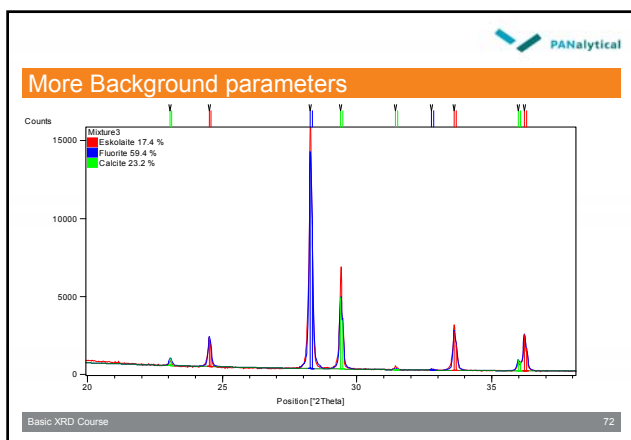
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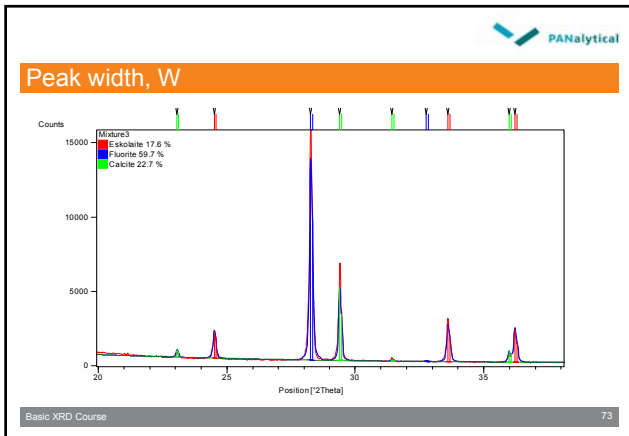
Identification with X'Pert HighScore: Search-match and Identification Background Information







Identification with X'Pert HighScore: Search-match and Identification Background Information



Amorphous Quantification(1)

Analyzing an amorphous material using the Rietveld calculations in HighScore Plus

1. An internal standard must be added to the sample in a known amount prior to the data collection.
2. Collect Rietveld quality data
3. Perform a Rietveld refinement.
4. Indicate the amount of the internal standard phase (Si).

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The slide contains a title 'Amorphous Quantification(1)' and a subtitle 'Analyzing an amorphous material using the Rietveld calculations in HighScore Plus'. Below the subtitle is a numbered list of four steps: 1. An internal standard must be added to the sample in a known amount prior to the data collection. 2. Collect Rietveld quality data. 3. Perform a Rietveld refinement. 4. Indicate the amount of the internal standard phase (Si). The slide footer includes 'Basic XRD Course' and the number '74'.

Amorphous Quantification(2)

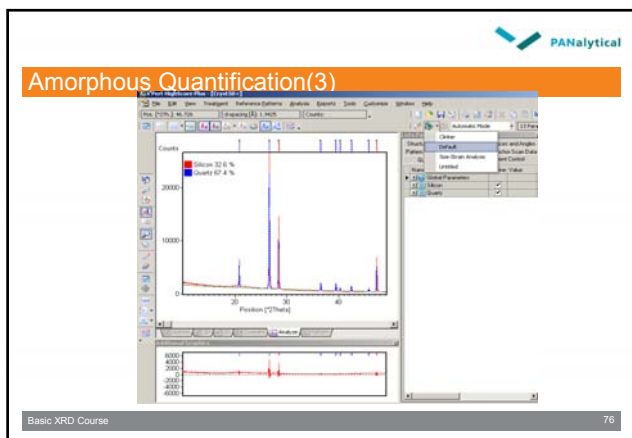
This example shows how to determine the amorphous content of a sample with a Rietveld refinement by addition of a crystalline standard phase.

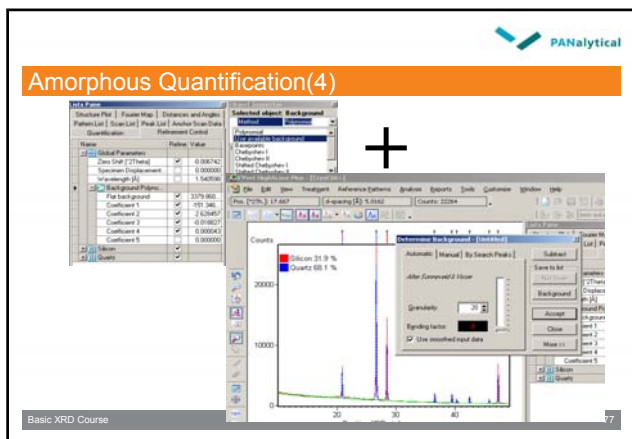
Amorphous compound(s) are invisible for the Rietveld method. Only crystalline phases are taken into account and their sum is normalized to 100%. The amount of the crystalline phases is overestimated in case amorphous material is present too.

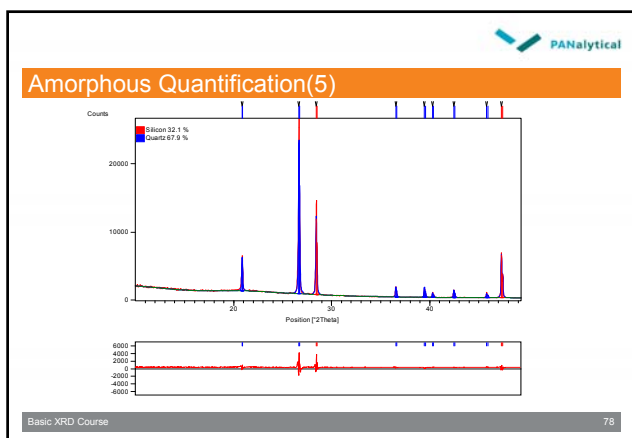
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The slide contains a title 'Amorphous Quantification(2)'. The main text explains that the Rietveld method is used to determine amorphous content by adding a crystalline standard phase. It notes that amorphous compounds are invisible to this method, and only crystalline phases are accounted for, with their sum normalized to 100%. This leads to an overestimation of crystalline phases when amorphous material is present. The slide footer includes 'Basic XRD Course' and the number '75'.

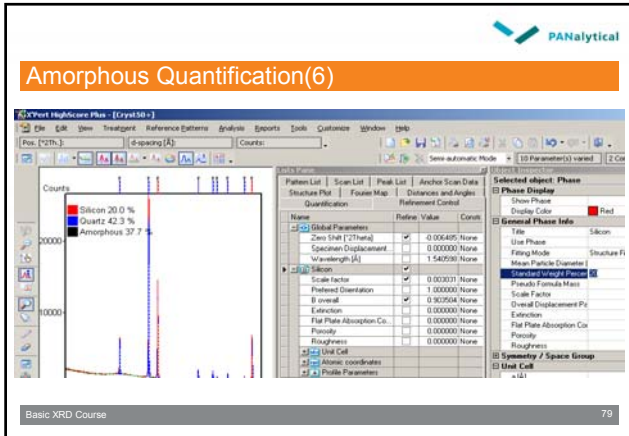
Identification with X'Pert HighScore: Search-match and Identification Background Information

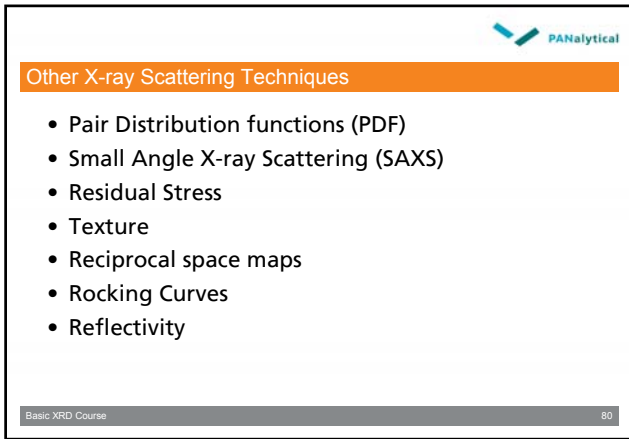


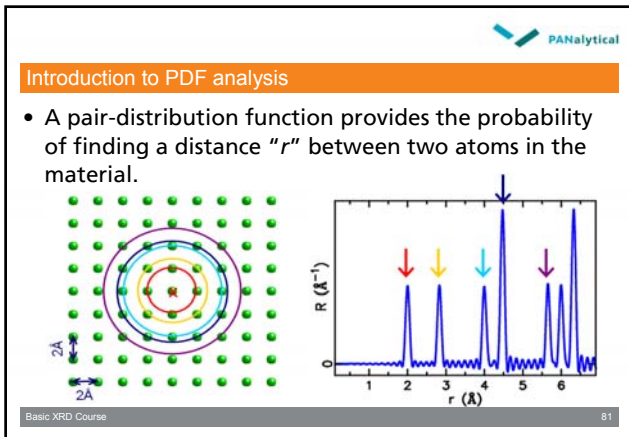




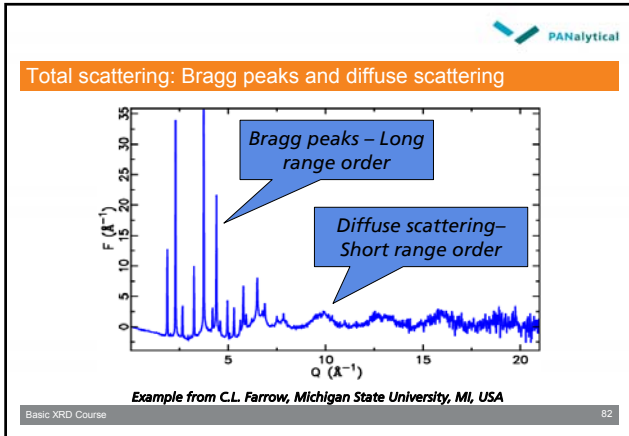
Identification with X'Pert HighScore: Search-match and Identification Background Information

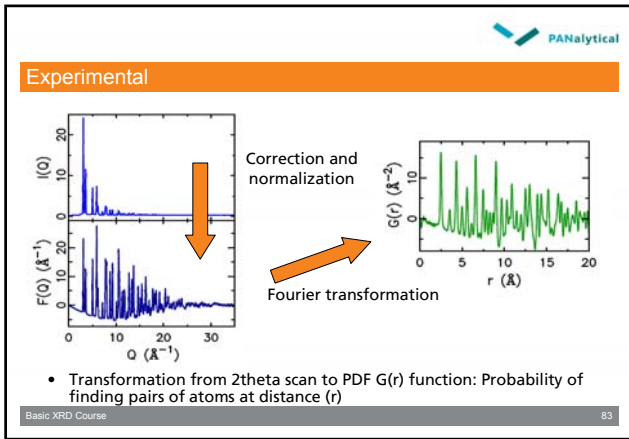


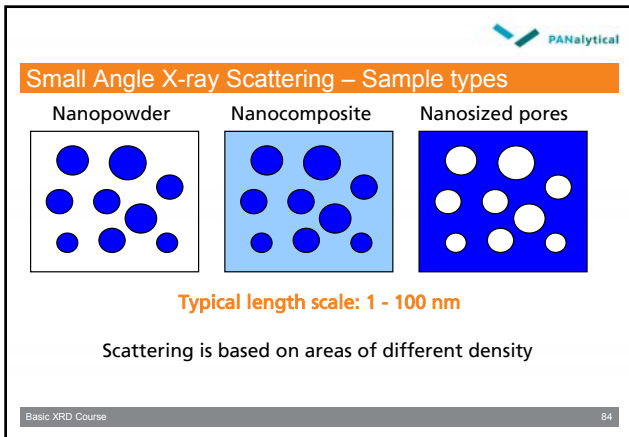




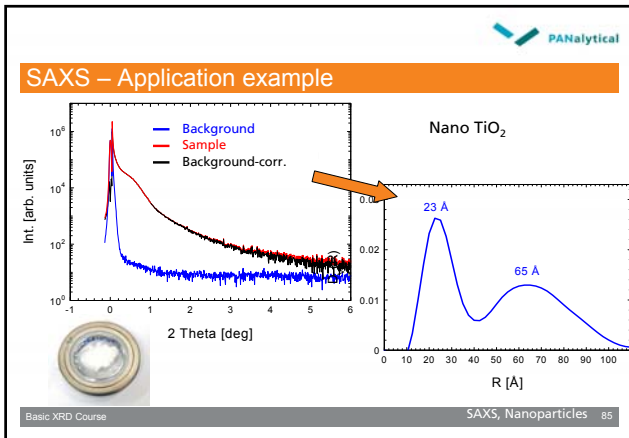
Identification with X'Pert HighScore: Search-match and Identification Background Information

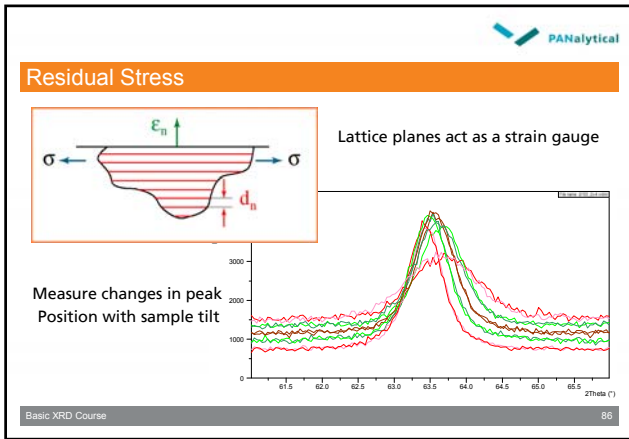


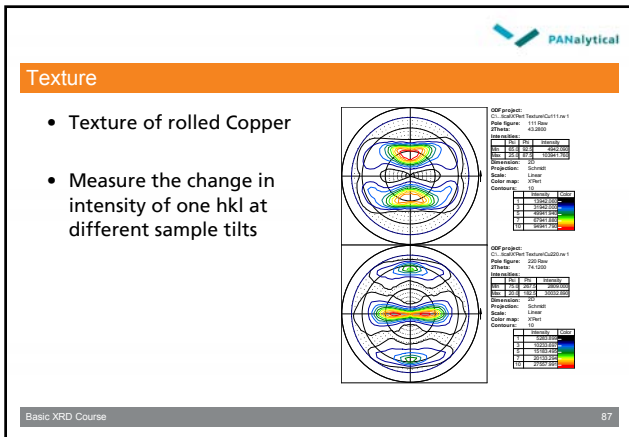





Identification with X'Pert HighScore: Search-match and Identification Background Information



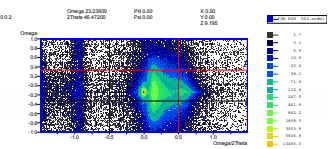




Identification with X'Pert HighScore: Search-match and Identification Background Information




Reciprocal space maps

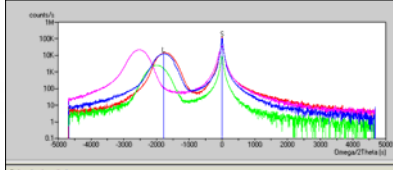


- Reciprocal space map of epitaxial SrMnO₂ on single crystal SrTiO₂
- Shows if the layer is strained or relaxed
- Shows the quality of the crystals

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Rocking Curves



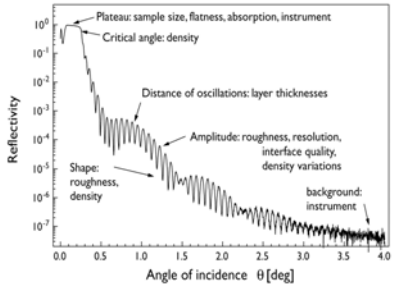
Refinement from single scans		Composition: Ga0.83009(6)160114u					
Bulk mismatch (ppm)	12125	Refinement (%)	88.9				
Parallel mismatch (ppm)	11765						
Calculate the fit & composition							
Scan	Date	hkl	Substrate	Substrate	Omega	Separation(s)	Phi(s)
Scan 1	Ga0.83009(6)160114u.d30	0 0 4	32.56216	65.16631	1745.9	90.00000	
Scan 2	Ga0.83009(6)160114u.d30	2 2 4	77.14601	83.47952	1999.4	90.00000	
Scan 3	Ga0.83009(6)160114u.d30	2 2 4	6.17703	83.95095	2537.5	90.00000	
Scan 4	Ga0.83009(6)160114u.d30						

Layer: 1: None, 2:nc, Bands: Peak = 0.100000 rad, Ga0.83009(6)160114u

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Reflectivity



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